

FIGURE 1A

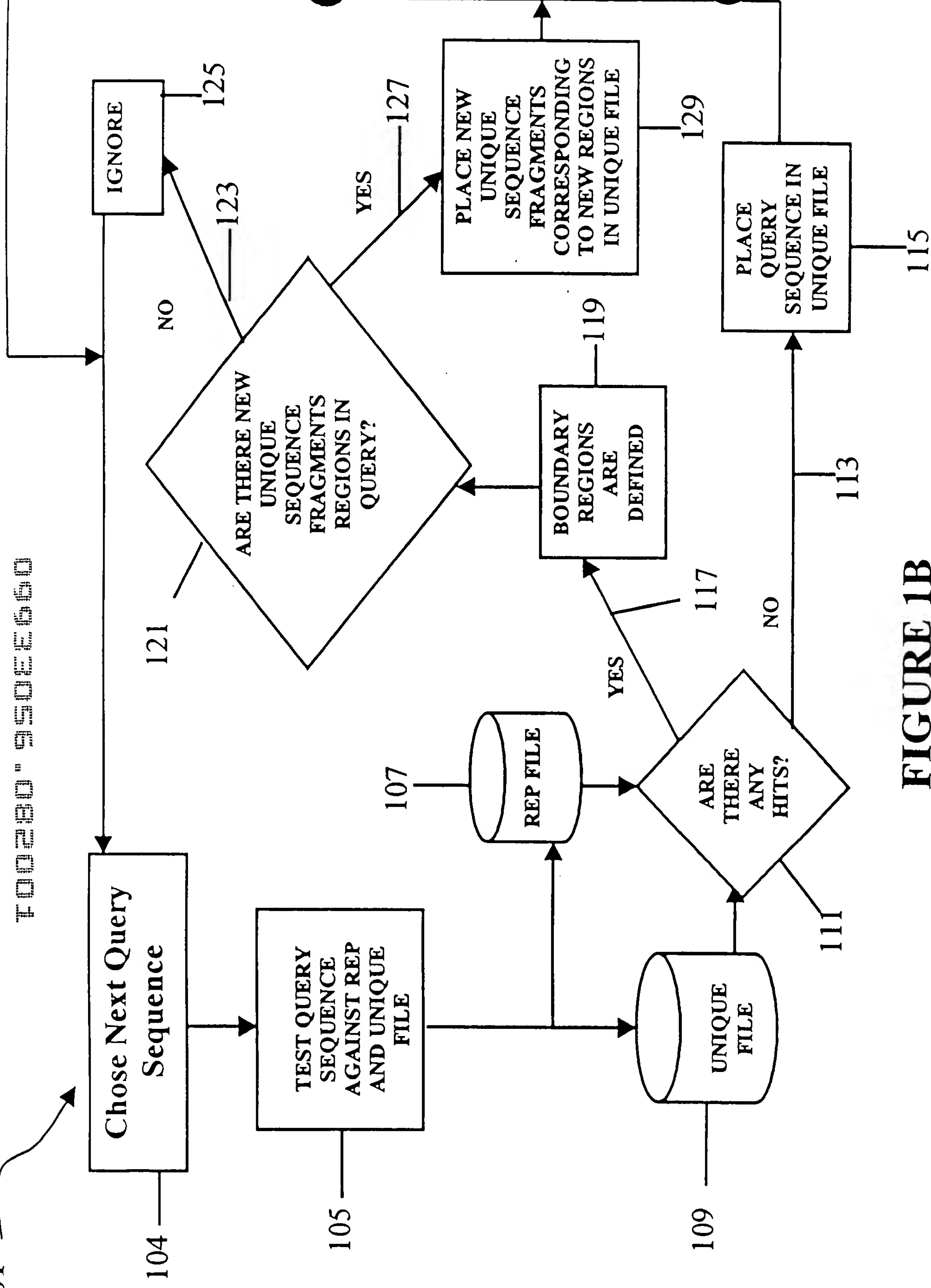


FIGURE 1B

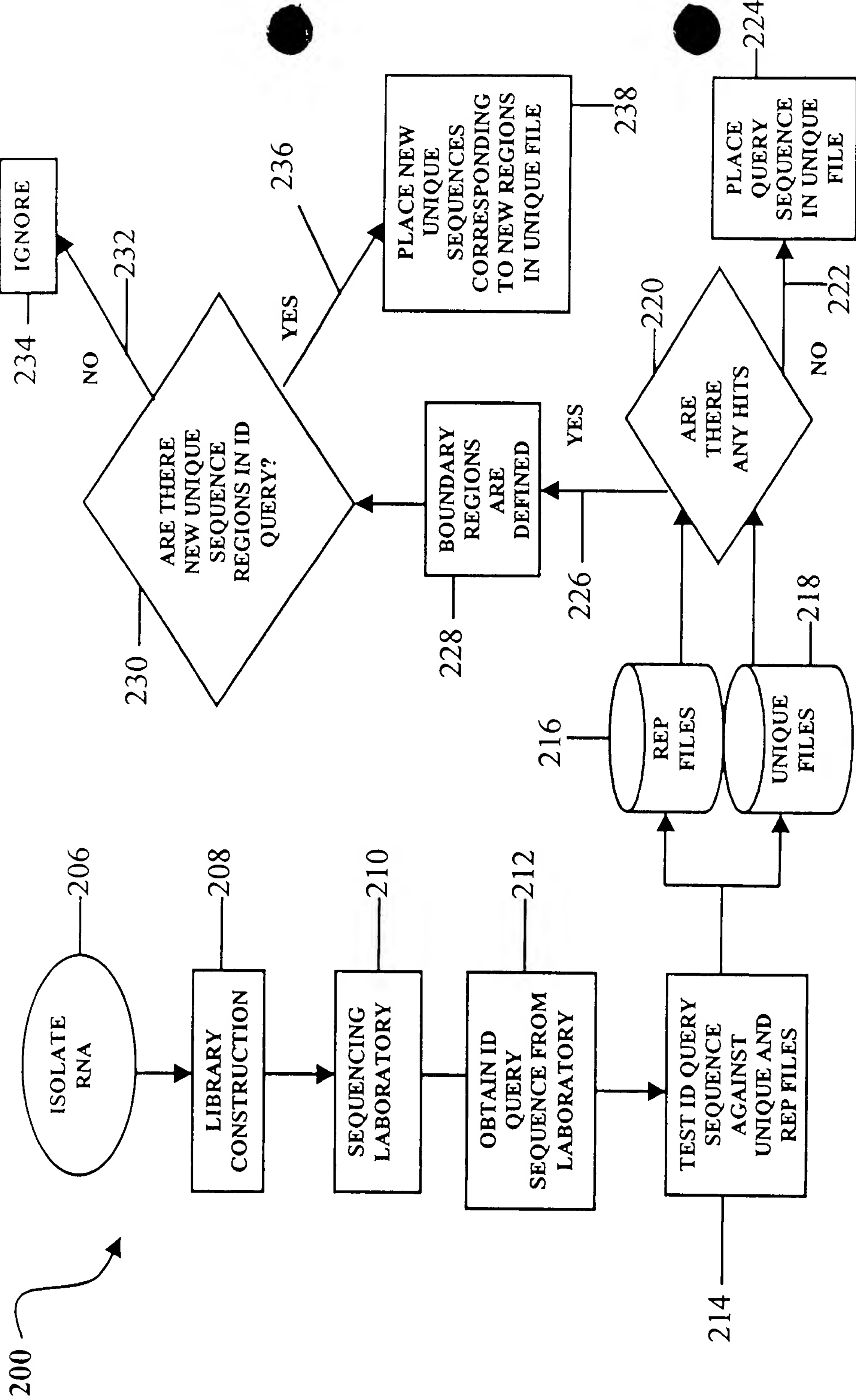
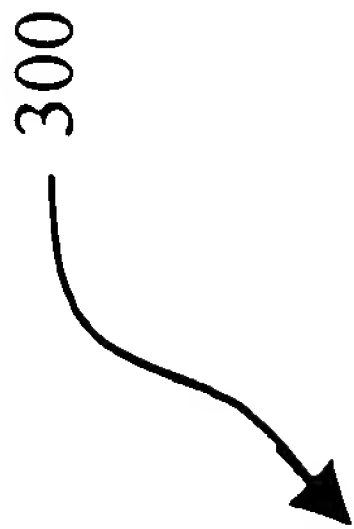


FIGURE 2

TCCGCGCA - AAACCTC



302

Query: Q_ileft

DBSeq: H_ileft

AATGGTGTAGGAGGG CGGCGCA -AAACCTC CANTCCCCCTGGGTTT AGG- -NTTTGGACCA AGTACCTCCAAATCG GGNGTTTGGCNCCA

AAAGGAGTA- GCGGG GGGTACTAACGGCTC CANTCCCCCTGGGTCT AGGATNTTTGGACCA CGCAC -TCCAAATCG GGNGTTTGGCNCCA



304

306

Q_iright

308

Figure 3

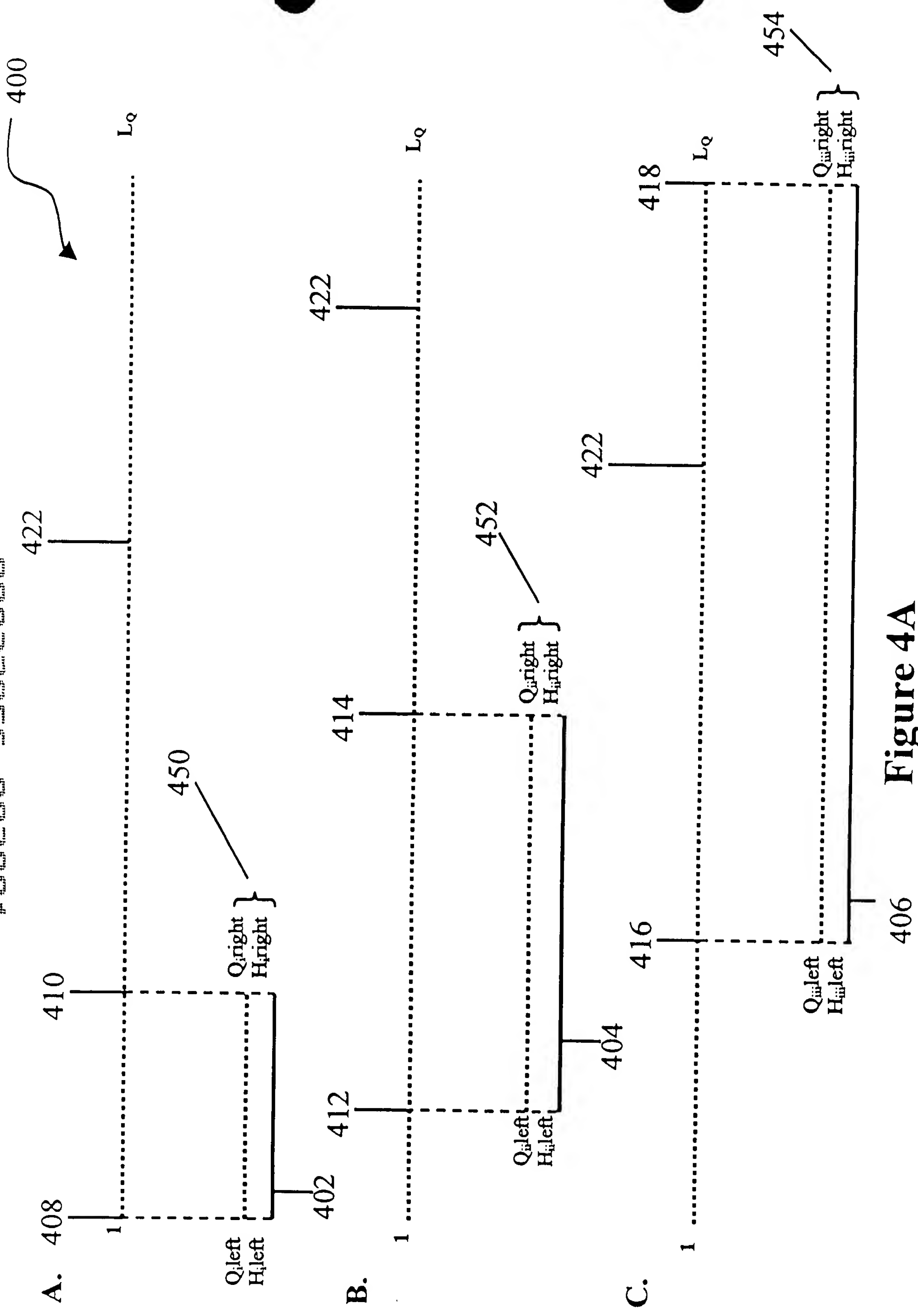


Figure 4A

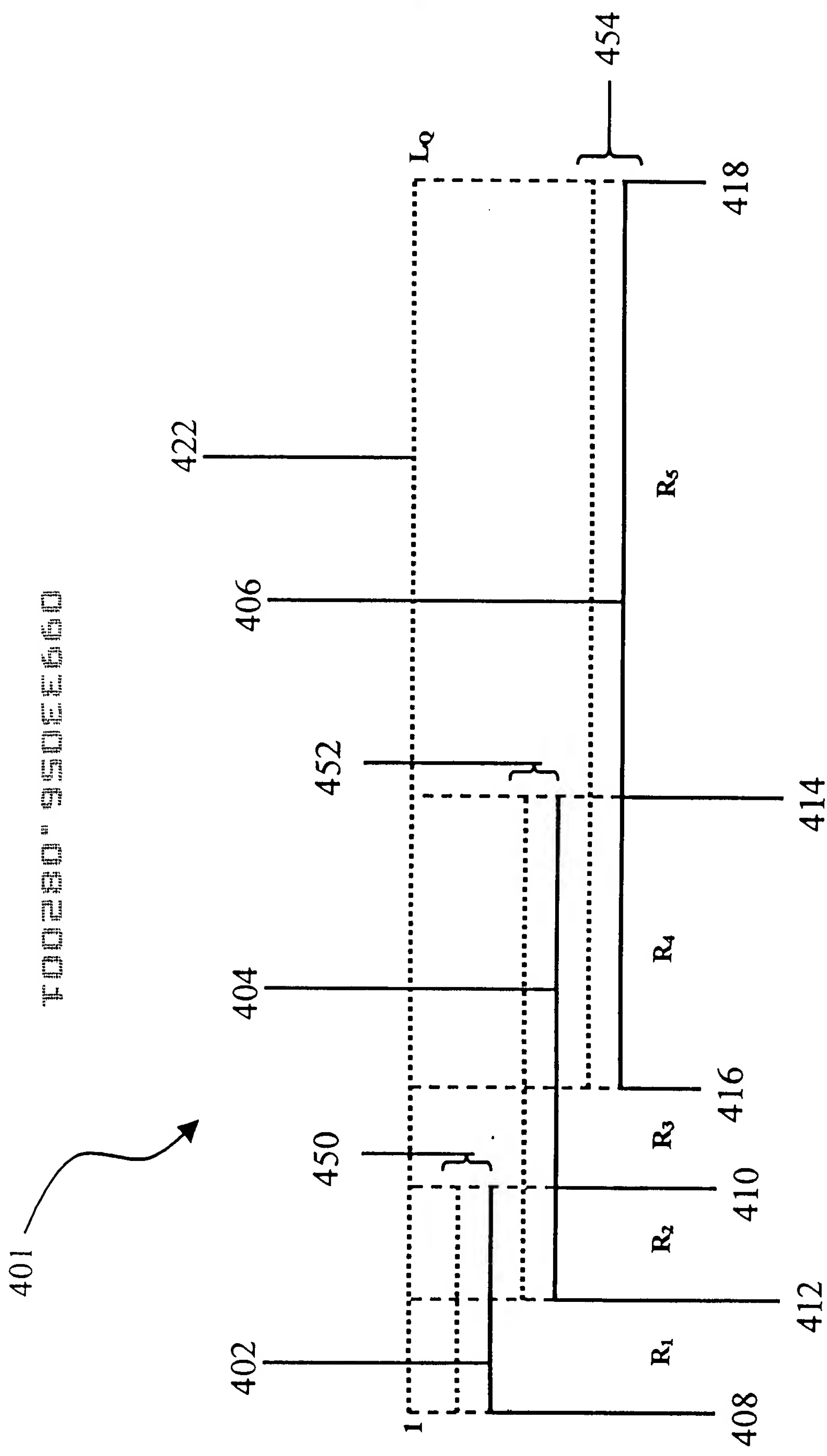


Figure 4B

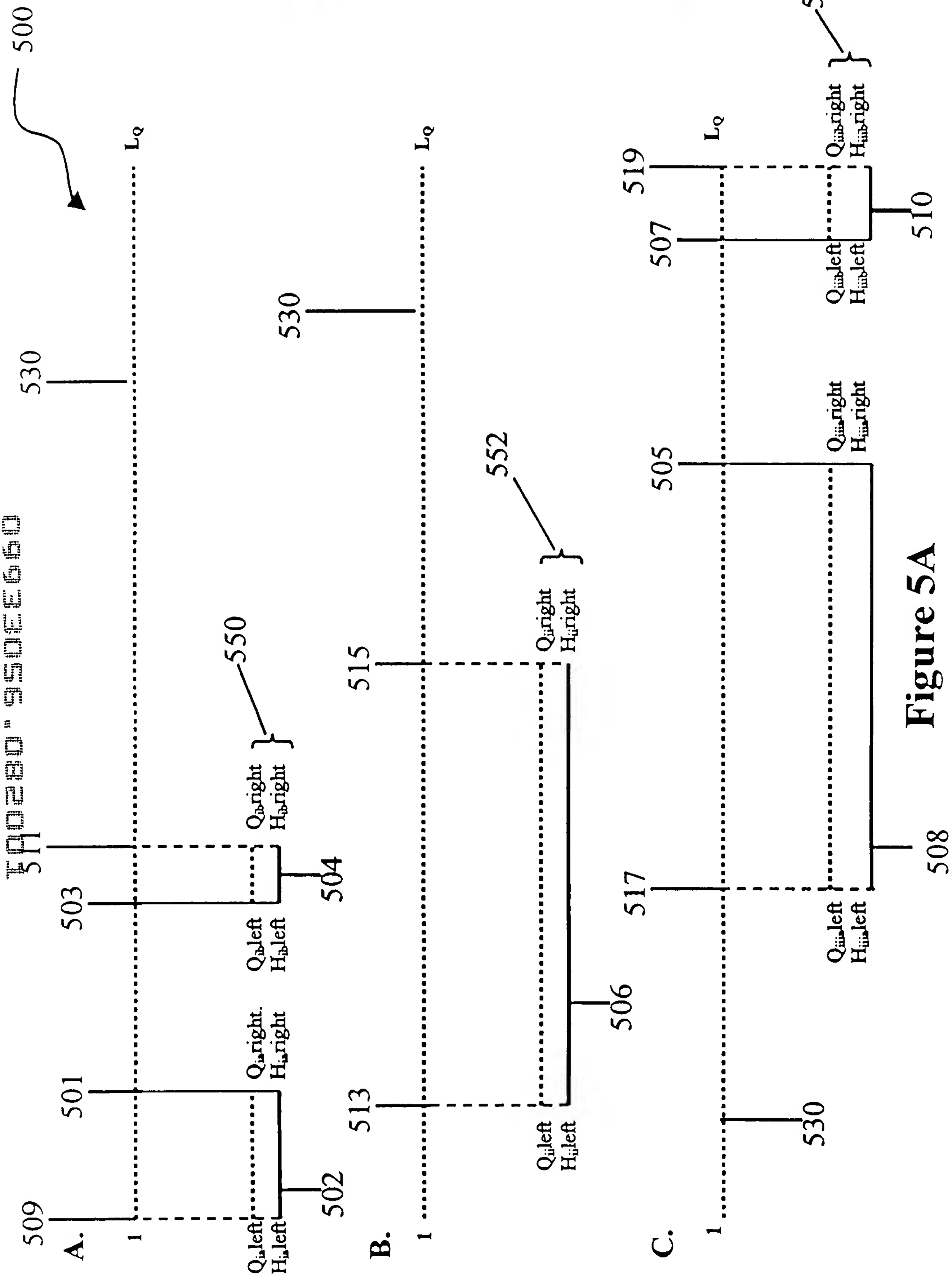


Figure 5A

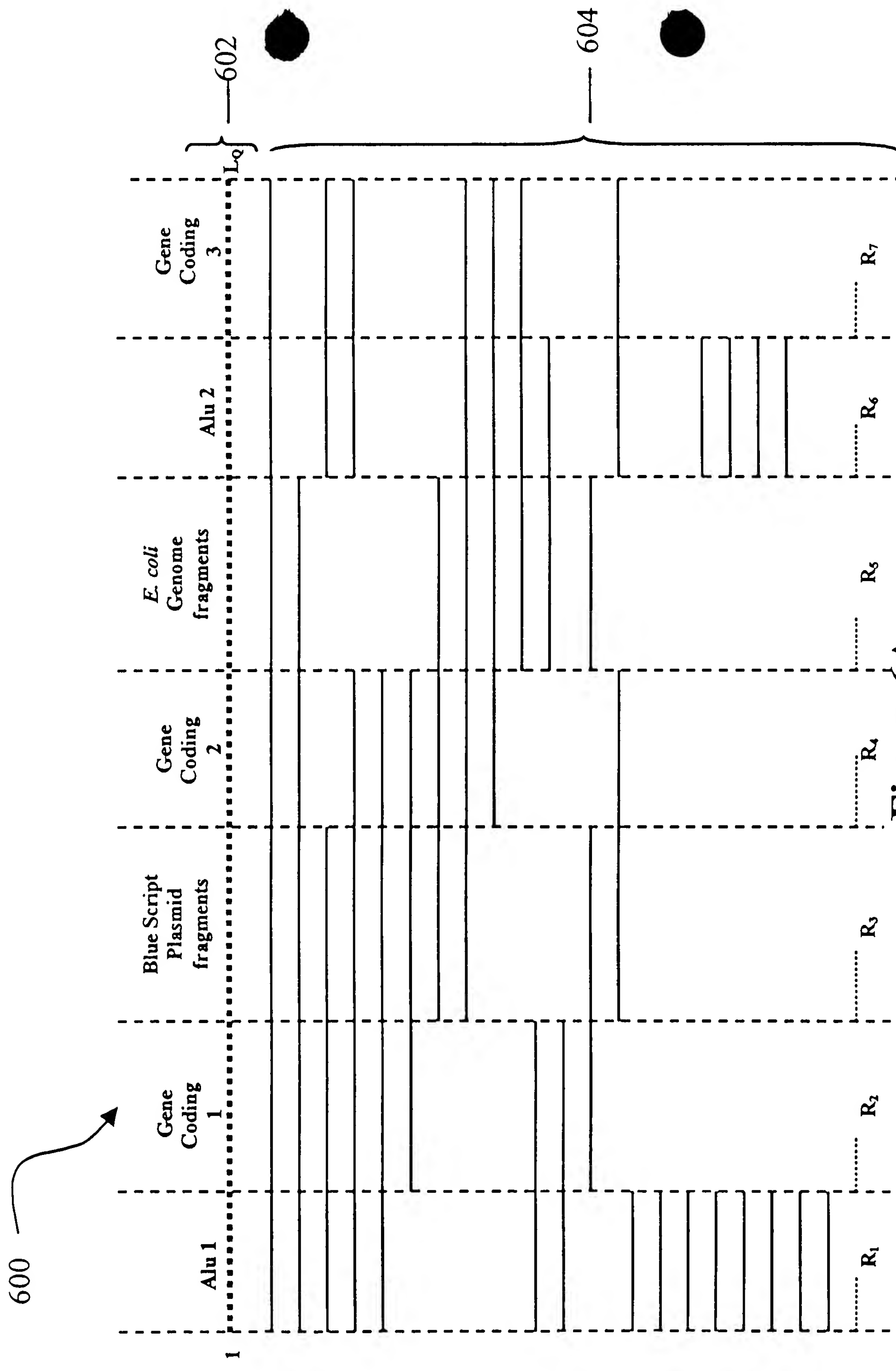
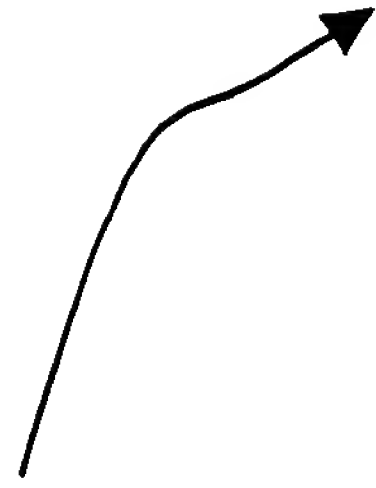


Figure 6A

601 

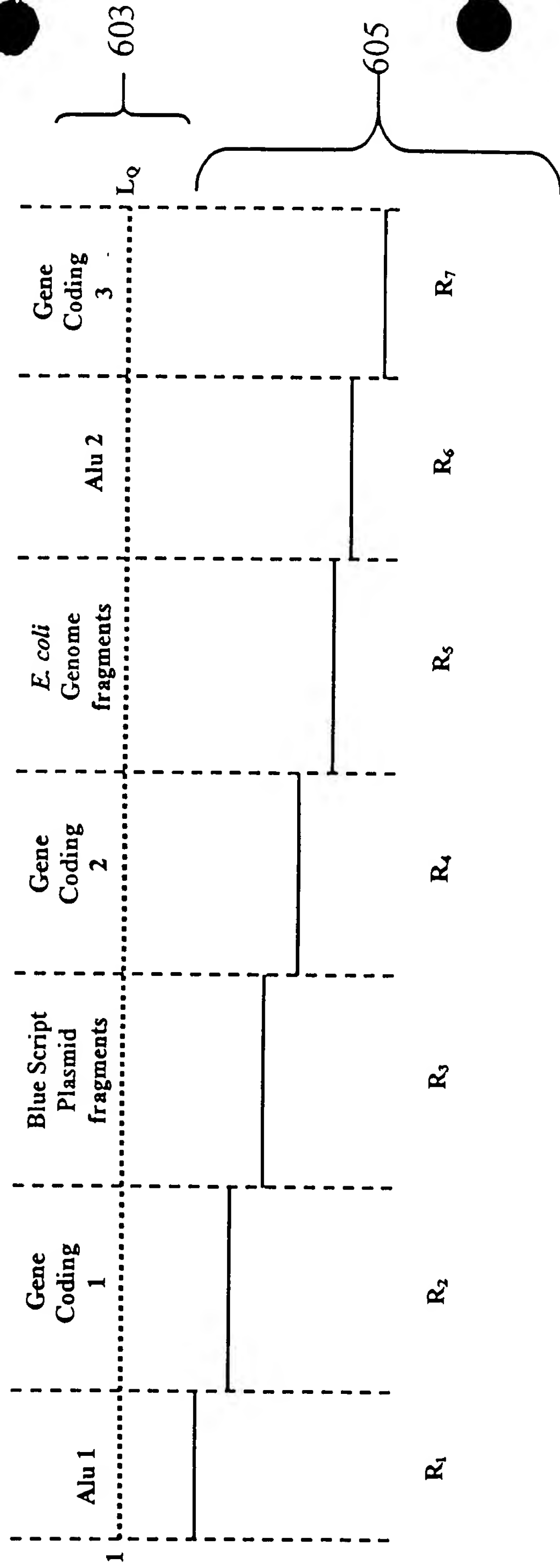


Figure 6B